SEQUENCE LISTING

<110> Fiedler, Dr. Ulrike
Rudolph, Prof. Dr. Rainer

<120> Design of beta-sheet proteins with specific binding properties

<130> P12389

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<151> 1999-07-13

<160> 22

<170> PatentIn Ver. 2.1

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<213> Artificial Sequence

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<210> 2
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: oligonucleotide

<400> 2
gccgcaggaa gtactggtga ccctggtagt tggggcgctc atacagcatc 50

<210> 3
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: oligonucleotide

<400> 3
ccatcagccc catcagcgaa ctttgccgca ggaagtactg g 41

<210> 4
<211> 48
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial

Sequence: oligonucleotide

<400> 4
gagtcattct gcggccgcat aaaaatccat cacccgtctt aaagaacc 48

<210> 5
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<223> Description of Artificial
Sequence: oligonucleotide

<400> 5
gcggcccagc cggccgctgc tggatgctgt atgagcgccc caactaccag ggtcaccag 59

<210> 6
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<212> DNA
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<220>
<223> Description of Artificial
Sequence: oligonucleotide

<400> 6
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<210> 7
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<213> Artificial Sequence

<220>
<223> Description of Artificial
Sequence: oligonucleotide

<400> 7
ccatgattac gccaagcttt ggagcc 26

<210> 8
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial
Sequence: oligonucleotide

<400> 8
ctgaaagtgc cggtgtgttg c 21

<210> 9
<211> 176
<212> DNA
<213> Bos sp.

<400> 9
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ttacagttgc aatagcgact gccccaacct gcagccctat ttcagccgct gtaactccat 120
cagggtgctg agcggctgct ggatgctgta tgagcgcccc aactaccagg gtcacc 176

<210> 10
<211> 176
<212> DNA
<213> Bos sp.

<400> 10
ggcccagccg gccatgggga agatcacttt ttacgaggac cggggcttcc agggccactg 60
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ccgcgtggac agcggctgct ggatgctgta tgagcgcccc aactaccagg gccacc 176

<210> 11
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<212> DNA
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<220>
<223> Description of Artificial Sequence: oligonucleotide

<400> 11
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<210> 12
<211> 36
<212> DNA
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<220>
<223> Description of Artificial Sequence: oligonucleotide

<400> 12
ggccatgggg nnkatcnnkt ttnnkgagga ccgggg 36

<210> 13
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
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<400> 13
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<210> 14
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: oligonucleotide

<400> 14
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<210> 15
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<400> 15
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<210> 16
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<212> DNA
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Sequence: oligonucleotide

<400> 16
gatggagtta cagcggctga aatagggctg caggttgggg cagtcgc 47

<210> 17
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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Sequence: oligonucleotide

<400> 17
tgtaactcca tcnnkgtgnn kagcggctgc tggatgctgt atgag 45

<210> 18
<211> 37
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial
Sequence: oligonucleotide

<400> 18
cgccccaact accagggtca ccagtacttc ctgcggc 37

<210> 19
<211> 198
<212> PRT
<213> Bos sp.

<400> 19
Ala Ala Gln Pro Ala Met Gly Arg Ile Lys Phe Lys Glu Asp Arg Gly
1 5 10 15

Phe Gln Gly His Tyr Tyr Ser Cys Asn Ser Asp Cys Pro Asn Leu Gln
20 25 30

Pro Tyr Phe Ser Arg Cys Asn Ser Ile Arg Val Leu Ser Gly Cys Trp
35 40 45

Met Leu Tyr Glu Arg Pro Asn Tyr Gln Gly His Gln Tyr Phe Leu Arg
50 55 60

Arg Gly Asp Tyr Pro Asp Tyr Gln Gln Trp Met Gly Phe Asn Asp Ser
65 70 75 80

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Ile Arg Ser Cys Arg Leu Ile Pro Gln His Thr Gly Thr Phe Arg Met
                85                  90                  95

Arg Ile Tyr Glu Arg Asp Asp Phe Arg Gly Gln Met Ser Glu Ile Thr
            100                 105                 110

Asp Asp Cys Pro Ser Leu Gln Asp Arg Phe His Leu Thr Glu Val His
        115                 120                 125

Ser Leu Asn Val Leu Glu Gly Ser Trp Val Leu Tyr Glu Met Pro Ser
    130                 135                 140

Tyr Arg Gly Arg Gln Tyr Leu Leu Arg Pro Gly Glu Tyr Arg Arg Tyr
145                 150                 155                 160

Leu Asp Trp Gly Ala Met Asn Ala Lys Val Gly Ser Leu Arg Arg Val
                165                 170                 175

Met Asp Phe Tyr Ala Ala Ala Gly Ala Pro Val Pro Tyr Pro Asp Pro
            180                 185                 190

Leu Glu Pro Arg Ala Ala
        195


<210> 20
<211> 198
<212> PRT
<213> Bos sp.

<400> 20
Ala Ala Gln Pro Ala Met Gly Lys Ile Thr Phe Tyr Glu Asp Arg Gly
  1               5                  10                  15

Phe Gln Gly His Cys Tyr Glu Cys Ser Ser Asp Cys Pro Asn Leu Gln
            20                  25                  30

Pro Tyr Phe Ser Arg Cys Asn Ser Ile Arg Val Asp Ser Gly Cys Trp
        35                  40                  45

Met Leu Tyr Glu Arg Pro Asn Tyr Gln Gly His Gln Tyr Phe Leu Arg
    50                  55                  60

Arg Gly Asp Tyr Pro Asp Tyr Gln Gln Trp Met Gly Phe Asn Asp Ser
65                  70                  75                  80

Ile Arg Ser Cys Arg Leu Ile Pro Gln His Thr Gly Thr Phe Arg Met
                85                  90                  95

Arg Ile Tyr Glu Arg Asp Asp Phe Arg Gly Gln Met Ser Glu Ile Thr
            100                 105                 110

Asp Asp Cys Pro Ser Leu Gln Asp Arg Phe His Leu Thr Glu Val His
        115                 120                 125

Ser Leu Asn Val Leu Glu Gly Ser Trp Val Leu Tyr Glu Met Pro Ser
    130                 135                 140

Tyr Arg Gly Arg Gln Tyr Leu Leu Arg Pro Gly Glu Tyr Arg Arg Tyr
145                 150                 155                 160

Leu Asp Trp Gly Ala Met Asn Ala Lys Val Gly Ser Leu Arg Arg Val
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165 170 175

Met Asp Phe Tyr Ala Ala Ala Gly Ala Pro Val Pro Tyr Pro Asp Pro
180 185 190

Leu Glu Pro Arg Ala Ala
195

<210> 21
<211> 197
<212> PRT
<213> Bos sp.

<400> 21
Met Gly Arg Ile Lys Phe Lys Glu Asp Arg Gly Phe Gln Gly His Tyr
1 5 10 15

Tyr Ser Cys Asn Ser Asp Cys Pro Asn Leu Gln Pro Tyr Phe Ser Arg
20 25 30

Cys Asn Ser Ile Arg Val Leu Ser Gly Cys Trp Met Leu Tyr Glu Arg
35 40 45

Pro Asn Tyr Gln Gly His Gln Tyr Phe Leu Arg Arg Gly Asp Tyr Pro
50 55 60

Asp Tyr Gln Gln Trp Met Gly Phe Asn Asp Ser Ile Arg Ser Cys Arg
65 70 75 80

Leu Ile Pro Gln His Thr Gly Thr Phe Arg Met Arg Ile Tyr Glu Arg
85 90 95

Asp Asp Phe Arg Gly Gln Met Ser Glu Ile Thr Asp Asp Cys Pro Ser
100 105 110

Leu Gln Asp Arg Phe His Leu Thr Glu Val His Ser Leu Asn Val Leu
115 120 125

Glu Gly Ser Trp Val Leu Tyr Glu Met Pro Ser Tyr Arg Gly Arg Gln
130 135 140

Tyr Leu Leu Arg Pro Gly Glu Tyr Arg Arg Tyr Leu Asp Trp Gly Ala
145 150 155 160

Met Asn Ala Lys Val Gly Ser Leu Arg Arg Val Met Asp Phe Tyr Ser
165 170 175

Asp Pro Asn Ser Ser Ser Val Asp Lys Leu Ala Ala Ala Leu Glu His
180 185 190

His His His His His
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<210> 22
<211> 197
<212> PRT
<213> Bos sp.

<400> 22
Met Gly Lys Ile Thr Phe Tyr Glu Asp Arg Gly Phe Gln Gly His Cys

P12389.sequence engl.txt

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  1               5                  10                  15

Tyr Glu Cys Ser Ser Asp Cys Pro Asn Leu Gln Pro Tyr Phe Ser Arg
            20                  25                  30

Cys Asn Ser Ile Arg Val Asp Ser Gly Cys Trp Met Leu Tyr Glu Arg
        35                  40                  45

Pro Asn Tyr Gln Gly His Gln Tyr Phe Leu Arg Arg Gly Asp Tyr Pro
    50                  55                  60

Asp Tyr Gln Gln Trp Met Gly Phe Asn Asp Ser Ile Arg Ser Cys Arg
65                  70                  75                  80

Leu Ile Pro Gln His Thr Gly Thr Phe Arg Met Arg Ile Tyr Glu Arg
                85                  90                  95

Asp Asp Phe Arg Gly Gln Met Ser Glu Ile Thr Asp Asp Cys Pro Ser
            100                 105                 110

Leu Gln Asp Arg Phe His Leu Thr Glu Val His Ser Leu Asn Val Leu
        115                 120                 125

Glu Gly Ser Trp Val Leu Tyr Glu Met Pro Ser Tyr Arg Gly Arg Gln
    130                 135                 140

Tyr Leu Leu Arg Pro Gly Glu Tyr Arg Arg Tyr Leu Asp Trp Gly Ala
145                 150                 155                 160

Met Asn Ala Lys Val Gly Ser Leu Arg Arg Val Met Asp Phe Tyr Ser
                165                 170                 175

Asp Pro Asn Ser Ser Ser Val Asp Lys Leu Ala Ala Ala Leu Glu His
            180                 185                 190

His His His His His
        195
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P12389.sequence engl.txt

P12389.sequence engl.txt